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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/859,604

DATE: 10/11/2001  
TIME: 09:21:13

Input Set : A:\Rih32ci1.app  
Output Set: N:\CRF3\10112001\I859604.raw

3 <110> APPLICANT: Wands, Jack R.  
 4 de la Monte, Suzanne M  
 5 Deutch, Alan H  
 6 Ghanbari, Hossein A  
 8 <120> TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS  
 10 <130> FILE REFERENCE: 21486-032 CIP  
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/859,604  
 13 <141> CURRENT FILING DATE: 2001-05-17  
 15 <150> PRIOR APPLICATION NUMBER: 09/436,184  
 16 <151> PRIOR FILING DATE: 1999-11-08  
 18 <160> NUMBER OF SEQ ID NOS: 13  
 20 <170> SOFTWARE: PatentIn Ver. 2.1  
 22 <210> SEQ ID NO: 1  
 23 <211> LENGTH: 36  
 24 <212> TYPE: PRT  
 25 <213> ORGANISM: Artificial Sequence  
 27 <220> FEATURE:  
 28 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus  
 29 EGF-like domain  
 31 <220> FEATURE:  
 32 <221> NAME/KEY: VARIANT  
 33 <222> LOCATION: (2)..(8)  
 34 <223> OTHER INFORMATION: Wherein any Xaa may be any amino acid  
 36 <220> FEATURE:  
 37 <221> NAME/KEY: VARIANT  
 38 <222> LOCATION: (10)..(13)  
 39 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
 41 <220> FEATURE:  
 42 <221> NAME/KEY: VARIANT  
 43 <222> LOCATION: (15)..(24)  
 44 <223> OTHER INFORMATION: Wherein Xaa is anu amino acid.  
 46 <220> FEATURE:  
 47 <221> NAME/KEY: VARIANT  
 48 <222> LOCATION: (26)  
 49 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
 51 <220> FEATURE:  
 52 <221> NAME/KEY: VARIANT  
 53 <222> LOCATION: (28)..(35)  
 54 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
 56 <400> SEQUENCE: 1  
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 58 1 5 10 15  
 W--> 60 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Cys Xaa Xaa Xaa Xaa Xaa  
 61 20 25 30  
 W--> 63 Xaa Xaa Xaa Cys  
 64 35  
 67 <210> SEQ ID NO: 2

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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/859,604

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Input Set : A:\Rih32cil.app  
Output Set: N:\CRF3\10112001\I859604.raw

68 <211> LENGTH: 758  
69 <212> TYPE: PRT  
70 <213> ORGANISM: Homo sapiens  
72 <400> SEQUENCE: 2  
73 Met Ala Gln Arg Lys Asn Ala Lys Ser Ser Gly Asn Ser Ser Ser Ser  
74 1 5 10 15  
76 Gly Ser Gly Ser Gly Ser Thr Ser Ala Gly Ser Ser Ser Pro Gly Ala  
77 20 25 30  
79 Arg Arg Glu Thr Lys His Gly Gly His Lys Asn Gly Arg Lys Gly Gly  
80 35 40 45  
82 Leu Ser Gly Thr Ser Phe Phe Thr Trp Phe Met Val Ile Ala Leu Leu  
83 50 55 60  
85 Gly Val Trp Thr Ser Val Ala Val Val Trp Phe Asp Leu Val Asp Tyr  
86 65 70 75 80  
88 Glu Glu Val Leu Gly Lys Leu Gly Ile Tyr Asp Ala Asp Gly Asp Gly  
89 85 90 95  
91 Asp Phe Asp Val Asp Asp Ala Lys Val Leu Leu Gly Leu Lys Glu Arg  
92 100 105 110  
94 Ser Thr Ser Glu Pro Ala Val Pro Pro Glu Glu Ala Glu Pro His Thr  
95 115 120 125  
97 Glu Pro Glu Glu Gln Val Pro Val Glu Ala Glu Pro Gln Asn Ile Glu  
98 130 135 140  
100 Asp Glu Ala Lys Glu Gln Ile Gln Ser Leu Leu His Glu Met Val His  
101 145 150 155 160  
103 Ala Glu His Val Glu Gly Glu Asp Leu Gln Gln Glu Asp Gly Pro Thr  
104 165 170 175  
106 Gly Glu Pro Gln Gln Glu Asp Asp Glu Phe Leu Met Ala Thr Asp Val  
107 180 185 190  
109 Asp Asp Arg Phe Glu Thr Leu Glu Pro Glu Val Ser His Glu Glu Thr  
110 195 200 205  
112 Glu His Ser Tyr His Val Glu Glu Thr Val Ser Gln Asp Cys Asn Gln  
113 210 215 220  
115 Asp Met Glu Glu Met Met Ser Glu Gln Glu Asn Pro Asp Ser Ser Glu  
116 225 230 235 240  
118 Pro Val Val Glu Asp Glu Arg Leu His His Asp Thr Asp Asp Val Thr  
119 245 250 255  
121 Tyr Gln Val Tyr Glu Glu Gln Ala Val Tyr Glu Pro Leu Glu Asn Glu  
122 260 265 270  
124 Gly Ile Glu Ile Thr Glu Val Thr Ala Pro Pro Glu Asp Asn Pro Val  
125 275 280 285  
127 Glu Asp Ser Gln Val Ile Val Glu Glu Val Ser Ile Phe Pro Val Glu  
128 290 295 300  
130 Glu Gln Gln Glu Val Pro Pro Glu Thr Asn Arg Lys Thr Asp Asp Pro  
131 305 310 315 320  
133 Glu Gln Lys Ala Lys Val Lys Lys Lys Pro Lys Leu Leu Asn Lys  
134 325 330 335  
136 Phe Asp Lys Thr Ile Lys Ala Glu Leu Asp Ala Ala Glu Lys Leu Arg  
137 340 345 350  
139 Lys Arg Gly Lys Ile Glu Glu Ala Val Asn Ala Phe Lys Glu Leu Val

**RAW SEQUENCE LISTING**  
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**Input Set : A:\Rih32cil.app**  
**Output Set: N:\CRF3\10112001\I859604.raw**

140	355	360	365
142	Arg Lys Tyr Pro Gln Ser Pro Arg Ala Arg Tyr Gly Lys Ala Gln Cys		
143	370	375	380
145	Glu Asp Asp Leu Ala Glu Lys Arg Arg Ser Asn Glu Val Leu Arg Gly		
146	385	390	395
148	Ala Ile Glu Thr Tyr Gln Glu Val Ala Ser Leu Pro Asp Val Pro Ala		
149	405	410	415
151	Asp Leu Leu Lys Leu Ser Leu Lys Arg Arg Ser Asp Arg Gln Gln Phe		
152	420	425	430
154	Leu Gly His Met Arg Gly Ser Leu Leu Thr Leu Gln Arg Leu Val Gln		
155	435	440	445
157	Leu Phe Pro Asn Asp Thr Ser Leu Lys Asn Asp Leu Gly Val Gly Tyr		
158	450	455	460
160	Leu Leu Ile Gly Asp Asn Asp Asn Ala Lys Lys Val Tyr Glu Glu Val		
161	465	470	475
163	Leu Ser Val Thr Pro Asn Asp Gly Phe Ala Lys Val His Tyr Gly Phe		
164	485	490	495
166	Ile Leu Lys Ala Gln Asn Lys Ile Ala Glu Ser Ile Pro Tyr Leu Lys		
167	500	505	510
169	Glu Gly Ile Glu Ser Gly Asp Pro Gly Thr Asp Asp Gly Arg Phe Tyr		
170	515	520	525
172	Phe His Leu Gly Asp Ala Met Gln Arg Val Gly Asn Lys Glu Ala Tyr		
173	530	535	540
175	Lys Trp Tyr Glu Leu Gly His Lys Arg Gly His Phe Ala Ser Val Trp		
176	545	550	555
178	Gln Arg Ser Leu Tyr Asn Val Asn Gly Leu Lys Ala Gln Pro Trp Trp		
179	565	570	575
181	Thr Pro Lys Glu Thr Gly Tyr Thr Glu Leu Val Lys Ser Leu Glu Arg		
182	580	585	590
184	Asn Trp Lys Leu Ile Arg Asp Glu Gly Leu Ala Val Met Asp Lys Ala		
185	595	600	605
187	Lys Gly Leu Phe Leu Pro Glu Asp Glu Asn Leu Arg Glu Lys Gly Asp		
188	610	615	620
190	Trp Ser Gln Phe Thr Leu Trp Gln Gln Gly Arg Arg Asn Glu Asn Ala		
191	625	630	635
193	Cys Lys Gly Ala Pro Lys Thr Cys Thr Leu Leu Glu Lys Phe Pro Glu		
194	645	650	655
196	Thr Thr Gly Cys Arg Arg Gly Gln Ile Lys Tyr Ser Ile Met His Pro		
197	660	665	670
199	Gly Thr His Val Trp Pro His Thr Gly Pro Thr Asn Cys Arg Leu Arg		
200	675	680	685
202	Met His Leu Gly Leu Val Ile Pro Lys Glu Gly Cys Lys Ile Arg Cys		
203	690	695	700
205	Ala Asn Glu Thr Arg Thr Trp Glu Glu Gly Lys Val Leu Ile Phe Asp		
206	705	710	715
208	Asp Ser Phe Glu His Glu Val Trp Gln Asp Ala Ser Ser Phe Arg Leu		
209	725	730	735
211	Ile Phe Ile Val Asp Val Trp His Pro Glu Leu Thr Pro Gln Gln Arg		
212	740	745	750

RAW SEQUENCE LISTING  
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Input Set : A:\Rih32c11.app  
Output Set: N:\CRF3\10112001\I859604.raw

214 Arg Ser Leu Pro Ala Ile  
215 . 755  
218 <210> SEQ ID NO: 3  
219 <211> LENGTH: 2324  
220 <212> TYPE: DNA  
221 <213> ORGANISM: Homo sapiens  
223 <400> SEQUENCE: 3  
224 cggaccgtgc aatggcccaag cgtaagaatg ccaagagcag cgccaacacg agcagcagcg 60  
225 gctccggcag cggtagcacg agtgcgggca gcagcagccc cggggcccg agagagacaa 120  
226 agcatggagg acacaagaat gggagggaaag gcggactctc gggacttca ttcttcacgt 180  
227 gtttatggt gattgcattt ctgggcgtct ggacatctgt agctgtcggt tggttgatc 240  
228 ttgttgacta tgaggaagtt ctaggaaaac taggaatcta ttagtgcgtat ggtgatggag 300  
229 attttgatgt ggatgatgcc aaagttttat taggacttaa agagagatct acttcagagc 360  
230 cagcagtccc gccagaagag gctgagccac acactgagcc cgaggagcag gttcctgtgg 420  
231 aggccagaacc ccagaataatc gaagatgaag caaaagaaca aattcagtcc cttctccatg 480  
232 aaatggtaca cgcagaacat gttgagggag aagacttgc acaagaagat ggacccacag 540  
233 gagaaccaca acaaggaggat gatgagtttgc ttatggcgac ttagtgcgtat gatagattt 600  
234 agaccctgga acctgaagta tctcatgaag aaaccgagca tagttaccac gtggaaagaga 660  
235 cagtttcaca agactgtaat caggatattttt aagagatgtat gtctgagcag gaaaatccag 720  
236 attccagtga accagtagta gaagatgaaa gattgcacca tgatacagat gatgtaacat 780  
237 accaagtcta tgaggaacaa gcagttatgtt aacctctaga aatgaaggg atagaaatca 840  
238 cagaagtaac tgctccccctt gaggataatc ctgttagaaga ttccacagta attgttagaaag 900  
239 aagtaagcat ttttcctgtg gaagaacacg aggaagtacc accagaaaca aatagaaaaa 960  
240 cagatgatcc agaacaaaaaa gcaaaagtta agaaaaagaa gcctaaactt ttaaataaaat 1020  
241 ttgataagac tattaaagctt gaaacttgcgtt ctgcagaaaa actccgtaaa agggaaaaa 1080  
242 ttgaggaagc agtgaatgca tttaaagaac tagtacgca ataccctcag agtccacgag 1140  
243 caagatatgg gaaggcgcag tgtgaggatg atttggctga gaagaggaga agtaatgagg 1200  
244 tgctacgtgg agccatcgag acgttaccaag aggtggccag cctacctgtat gtcctgcag 1260  
245 acctgctgaa gctgagtttgc aagcgtcgtt cagacaggca acaatttcta ggtcatatga 1320  
246 gaggttccctt gcttaccctt cagagattttt ttcaactatt tcccaatgtat acttccttaa 1380  
247 aaaatgaccc ttgcgtggaa taccttttgc taggagataa tgacaatgca aagaaaagttt 1440  
248 atgaagaggt gctgagttgtt acacctaatttgc atggctttgc taaagtccat tatggcttca 1500  
249 tcctgaaggc acagaacaaa attgctgaga gcatccata tttaaaggaa ggaatagaat 1560  
250 ccggagatcc tggcactgtat gatgggagat ttatccatc cctggggat gccatgcaga 1620  
251 ggggtggaa caaagaggca tataagtgtt atgagcttgg gcacaagaga ggacactttg 1680  
252 catctgtctg gcaacgtca ctctacaatg tgaatggact gaaagcacag ccttgggtgg 1740  
253 ccccaaaaga aacgggttac acagagtttgc taaagtctttt agaaagaaac tggaagttaa 1800  
254 tccgagatgtt aggccttgca gtgtatggata aagccaaagg tctttccctg cctgaggatg 1860  
255 aaaacctgtt gggaaaaagg gactggagcc agttcacgtt gtggcagcaa ggaagaagaa 1920  
256 atgaaaatgtt ctgcaaaaggaa gctctttttt cctgtacctt actagaaaaat ttccccgaga 1980  
257 caacaggatg cagaagagga cagatcaat attccatcat gcacccccc actcacgtgt 2040  
258 ggccgcacac agggcccaaca aactgcagcc tccgaatgca cctgggcttgc gtgatccca 2100  
259 aggaaggctt gcaagatttgc tttttttttt gggggggggaa ggcaagggtgc 2160  
260 tcatcttgc tttttttttt gggggggggaa ggcaagggtgc 2220  
261 tattcatgtt ggtatgttgg catccggaaac tgacaccaca gcagagacgc agccttccag 2280  
262 caattttgttgc tttttttttt gggggggggaa aaactcttgc gaga 2324  
265 <210> SEQ ID NO: 4  
266 <211> LENGTH: 31  
267 <212> TYPE: PRT

RAW SEQUENCE LISTING  
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Input Set : A:\Rih32ci1.app  
Output Set: N:\CRF3\10112001\I859604.raw

268 <213> ORGANISM: Artificial Sequence  
270 <220> FEATURE:  
271 <223> OTHER INFORMATION: Description of Artificial Sequence: EGF-like  
272 cysteine-rich repeat  
274 <220> FEATURE:  
275 <221> NAME/KEY: VARIANT  
276 <222> LOCATION: (2)..(5)  
277 <223> OTHER INFORMATION: Wherein any Xaa may be any amino acid  
279 <220> FEATURE:  
280 <221> NAME/KEY: VARIANT  
281 <222> LOCATION: (7)..(8)  
282 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
284 <220> FEATURE:  
285 <221> NAME/KEY: VARIANT  
286 <222> LOCATION: (10)  
287 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
289 <220> FEATURE:  
290 <221> NAME/KEY: VARIANT  
291 <222> LOCATION: (14)  
292 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
294 <220> FEATURE:  
295 <221> NAME/KEY: VARIANT  
296 <222> LOCATION: (17)..(18)  
297 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
299 <220> FEATURE:  
300 <221> NAME/KEY: VARIANT  
301 <222> LOCATION: (25)..(26)  
302 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
304 <220> FEATURE:  
305 <221> NAME/KEY: VARIANT  
306 <222> LOCATION: (29)  
307 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.  
309 <400> SEQUENCE: /4 /  
W--> 310 Cys Asp Xaa Xaa Xaa Cys Xaa Xaa Lys Xaa Gly Asn Gly Xaa Cys Asp  
311 1 5 10 15  
W--> 313 Xaa Xaa Cys Asn Asn Ala Ala Cys Xaa Xaa Asp Gly Xaa Asp Cys  
314 20 25 30  
317 <210> SEQ ID NO: 5  
318 <211> LENGTH: 1242  
319 <212> TYPE: PRT  
320 <213> ORGANISM: Homo sapiens  
322 <400> SEQUENCE: 5  
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324 1 5 10 15  
326 Gly Tyr Leu Arg Lys Pro Lys Ser Met His Lys Arg Phe Phe Val Leu  
327 20 25 30  
329 Arg Ala Ala Ser Glu Ala Gly Gly Pro Ala Arg Leu Glu Tyr Tyr Glu  
330 35 40 45  
332 Asn Glu Lys Lys Trp Arg His Lys Ser Ser Ala Pro Lys Arg Ser Ile

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/09/859,604

DATE: 10/11/2001  
TIME: 09:21:14

Input Set : A:\Rih32cil.app  
Output Set: N:\CRF3\10112001\I859604.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:60 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:63 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:313 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4